

#2
OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/927,091

DATE: 08/17/2001
TIME: 12:03:45

Input Set : A:\Utsc651.app
Output Set: N:\CRF3\08162001\I927091.raw

3 <110> APPLICANT: KILLARY, ANN
 4 LOTT, STEVE
 5 CHANDLER, DAWN
 7 <120> TITLE OF INVENTION: THE TUMOR SUPPRESSOR CAR-1
 9 <130> FILE REFERENCE: UTSC:651US
 OC--> 11 <140> CURRENT APPLICATION NUMBER: US/09/927,091
 12 <141> CURRENT FILING DATE: 2001-08-09
 14 <150> PRIOR APPLICATION NUMBER: 60/227,560
 15 <151> PRIOR FILING DATE: 2000-08-23
 17 <150> PRIOR APPLICATION NUMBER: 60/225,033
 18 <151> PRIOR FILING DATE: 2000-08-10
 20 <160> NUMBER OF SEQ ID NOS: 9
 22 <170> SOFTWARE: PatentIn Ver. 2.1
 24 <210> SEQ ID NO: 1
 25 <211> LENGTH: 475
 26 <212> TYPE: PRT
 27 <213> ORGANISM: Human
 29 <400> SEQUENCE: 1
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 31 1 5 10 15
 33 Ile Tyr Gln Asp Pro Val Ser Leu Gly Cys Glu His Tyr Phe Cys Arg
 34 20 25 30
 36 Arg Cys Ile Thr Glu His Trp Val Arg Gln Glu Ala Gln Gly Ala Arg
 37 35 40 45
 39 Asp Cys Pro Glu Cys Arg Arg Thr Phe Ala Glu Pro Ala Leu Ala Pro
 40 50 55 60
 42 Ser Leu Lys Leu Ala Asn Ile Val Glu Arg Tyr Ser Ser Phe Pro Leu
 43 65 70 75 80
 45 Asp Ala Ile Leu Asn Ala Arg Arg Ala Ala Arg Pro Cys Gln Ala His
 46 85 90 95
 48 Asp Lys Val Lys Leu Phe Cys Leu Thr Asp Arg Ala Leu Leu Cys Phe
 49 100 105 110
 51 Phe Cys Asp Glu Pro Ala Leu His Glu Gln His Gln Val Thr Gly Ile
 52 115 120 125
 54 Asp Asp Ala Phe Asp Glu Leu Gln Arg Glu Leu Lys Asp Gln Leu Gln
 55 130 135 140
 57 Ala Leu Gln Asp Ser Glu Arg Glu His Thr Glu Ala Leu Gln Leu Leu
 58 145 150 155 160
 60 Lys Arg Gln Leu Ala Glu Thr Lys Ser Ser Thr Lys Ser Leu Arg Thr
 61 165 170 175
 63 Thr Ile Gly Glu Ala Phe Glu Arg Leu His Arg Leu Leu Arg Glu Arg
 64 180 185 190
 66 Gln Lys Ala Met Leu Glu Glu Leu Glu Ala Asp Thr Ala Arg Thr Leu
 67 195 200 205
 69 Thr Asp Ile Glu Gln Lys Val Gln Arg Tyr Ser Gln Gln Leu Arg Lys
 70 210 215 220
 72 Val Gln Glu Gly Ala Gln Ile Leu Gln Glu Arg Leu Ala Glu Thr Asp

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73	225	230	235	240
75	Arg His Thr Phe Leu Ala Gly Val Ala Ser Leu Ser Glu Arg Leu Lys			
76	245	250	255	
78	Gly Lys Ile His Glu Thr Asn Leu Thr Tyr Glu Asp Phe Pro Thr Ser			
79	260	265	270	
81	Lys Tyr Thr Gly Pro Leu Gln Tyr Thr Ile Trp Lys Ser Leu Phe Gln			
82	275	280	285	
84	Asp Ile His Pro Val Pro Ala Ala Leu Thr Leu Asp Pro Gly Thr Ala			
85	290	295	300	
87	His Gln Arg Leu Ile Leu Ser Asp Asp Cys Thr Ile Val Ala Tyr Gly			
88	305	310	315	320
90	Asn Leu His Pro Gln Pro Leu Gln Asp Ser Pro Lys Arg Phe Asp Val			
91	325	330	335	
93	Glu Val Ser Val Leu Gly Ser Glu Ala Phe Ser Ser Gly Val His Tyr			
94	340	345	350	
96	Trp Glu Val Val Val Ala Glu Lys Thr Gln Trp Val Ile Gly Leu Ala			
97	355	360	365	
99	His Glu Ala Ala Ser Arg Lys Gly Ser Ile Gln Ile Gln Pro Ser Arg			
100	370	375	380	
102	Gly Phe Tyr Cys Ile Val Met His Asp Gly Asn Gln Tyr Ser Ala Cys			
103	385	390	395	400
105	Thr Glu Pro Trp Thr Arg Leu Asn Val Arg Asp Lys Leu Asp Lys Val			
106	405	410	415	
108	Gly Val Phe Leu Asp Tyr Asp Gln Gly Leu Leu Ile Phe Tyr Asn Ala			
109	420	425	430	
111	Asp Asp Met Ser Trp Leu Tyr Thr Phe Arg Glu Lys Phe Pro Gly Lys			
112	435	440	445	
114	Leu Cys Ser Tyr Phe Ser Pro Gly Gln Ser His Ala Asn Gly Lys Asn			
115	450	455	460	
117	Val Gln Pro Leu Arg Ile Asn Thr Val Arg Ile			
118	465	470	475	
121	<210> SEQ ID NO: 2			
122	<211> LENGTH: 304			
123	<212> TYPE: PRT			
124	<213> ORGANISM: Human			
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130	Ile Tyr Gln Asp Pro Val Ser Leu Gly Cys Glu His Tyr Phe Cys Arg			
131	20	25	30	
133	Arg Cys Ile Thr Glu His Trp Val Arg Gln Glu Ala Gln Gly Ala Arg			
134	35	40	45	
136	Asp Cys Pro Glu Cys Arg Arg Thr Phe Ala Glu Pro Ala Leu Ala Pro			
137	50	55	60	
139	Ser Leu Lys Leu Ala Asn Ile Val Glu Arg Tyr Ser Ser Phe Pro Leu			
140	65	70	75	80
142	Asp Ala Ile Leu Asn Ala Arg Arg Ala Ala Arg Pro Cys Gln Ala His			
143	85	90	95	
145	Asp Lys Val Lys Leu Phe Cys Leu Thr Asp Arg Ala Leu Leu Cys Phe			

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146	100	105	110
148	Phe Cys Asp Glu Pro Ala Leu His Glu Gln His Gln Val Thr Gly Ile		
149	115	120	125
151	Asp Asp Ala Phe Asp Glu Leu Gln Arg Glu Leu Lys Asp Gln Leu Gln		
152	130	135	140
154	Ala Leu Gln Asp Ser Glu Arg Glu His Thr Glu Ala Leu Gln Leu Leu		
155	145	150	155
156	Lys Arg Gln Leu Ala Glu Thr Lys Ser Ser Thr Lys Ser Leu Arg Thr		
158	165	170	175
160	Thr Ile Gly Glu Ala Phe Glu Arg Leu His Arg Leu Leu Arg Glu Arg		
161	180	185	190
163	Gln Lys Ala Met Leu Glu Glu Leu Glu Ala Asp Thr Ala Arg Thr Leu		
164	195	200	205
166	Thr Asp Ile Glu Gln Lys Val Gln Arg Tyr Ser Gln Gln Leu Arg Lys		
167	210	215	220
169	Val Gln Glu Gly Ala Gln Ile Leu Gln Glu Arg Leu Ala Glu Thr Asp		
170	225	230	235
172	Arg His Thr Phe Leu Ala Gly Val Ala Ser Leu Ser Glu Arg Ala Ser		
173	245	250	255
175	Arg Pro Asn Pro Gly Pro Gly His Ser Pro Pro Ala Pro Asp Pro Val		
176	260	265	270
178	Gly Arg Leu His His Cys Gly Leu Arg Gln Leu Ala Pro Thr Ala Thr		
179	275	280	285
181	Ala Gly Leu Ala Lys Ala Leu Arg Cys Gly Gly Val Gly Ala Gly Phe		
182	290	295	300
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189	<211> LENGTH: 3826		
190	<212> TYPE: DNA		
191	<213> ORGANISM: Human		
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196	gtcacagcc aatgtacggc tcggcctggc tgccccctcc cccaggattc cccatccccca 180		
197	gcttcgtccgc ctcggccac cggcccccacc cccggatttc gaccggctta agggctccac 240		
198	cccgctccgg gatccccctt tcccaagctcc tatcccttag gactgccccg ccccccataaa 300		
199	cctcccccgtc aggatctccg tccctcagcc gtcacagcc tcctcccaagc gccccatcgcc 360		
200	ttgagctgcc cactacctt agactgccct cccgggttgg cgtcccaaggg agtctcagcc 420		
201	gcgcacccct tcctcggtt accctcctt cggacagcac cccctccctt ctccggtagc 480		
202	tcctacccct gcctgtcgg gcctcggtcc cgcggccacg cctcggtgt gcctccgaca 540		
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204	gccatggcggt gcagcctcaa ggacgagctg ctgtgtccaa tctgccttag catctaccag 660		
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206	gtgcggcagg aggccgaggc cggccgcac tgccccgagt gccggcgcac gttcgccgag 780		
207	cccgcgctgg cggccagcct caagctggcc aacatcggtt agcgctacag ctccctcccg 840		
208	ctggacgcca tcctcaacgc gcggccgcgc ggcgcgaccct gccaggcgcac cgacaaggtc 900		
209	aagctttct gcctcacgga cccgcgcgtt ctctgtttt tctgcgtacga gcctgcactg 960		
210	cacgagcagc atcaggtcac cggcatcgac gacgccttcg acgagctgca gagggagctg 1020		
211	aaggaccaac ttccaggccct tcaagacagc gagcggaaac acacccgaagc gctgcagctg 1080		
212	ctcaagcgac aactggcgga gaccaagtct tccaccaaga gcctgcggac cactatcgcc 1140		

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/927,091

DATE: 08/17/2001

TIME: 12:03:46

Input Set : A:\Utsc651.app

Output Set: N:\CRF3\08162001\I927091.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:1188 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:1619 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:2490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:2491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:2492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:2796 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:2921 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:2922 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:2923 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3010 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3011 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3012 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3044 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3045 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3057 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3115 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3150 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3319 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3320 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:3321 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8